

2610-Pos**Development of an *in vivo* Model to Investigate the Role of the Heavy Metal Efflux System Sil from *Cupriavidus Metallidurans* Ch34 in Resistance to Silver and Copper Ions**Elisabeth Ngonlong Ekendé¹, Max Mergeay², Jean-Marie Ruysschaert¹, Guy Vandenbussche¹.¹Laboratory for the Structure and Function of Biological Membranes, Center for Structural Biology and Bioinformatics, Université Libre de Bruxelles, B-1050 Bruxelles, Belgium, ²Research Unit for Microbiology, Belgian Nuclear Research Centre, SCK•CEN, B-2400-Mol, Belgium.

The RND-type (Resistance Nodulation and cell Division) efflux systems play an important role in the ability of bacteria to survive in the presence of a broad range of toxic compounds. Two major subfamilies are represented by the systems involved in Heavy Metal Efflux (HME-RND) or Hydrophobic and Amphiphilic compounds (e.g. antibiotics) Efflux (HAE-RND). These tripartite protein complexes are composed of an inner membrane (RND) and an outer membrane (member of the Outer Membrane Factor family - OMF) component linked together by a periplasmic adaptor protein (member of the Membrane Fusion Protein family - MFP). The RND protein is a cation/proton antiporter and is responsible for the substrate specificity. SilABC is an uncharacterized HME-RND system from *C. metallidurans* CH34. Using a proteomic approach, we have previously demonstrated the induction of Sil proteins in response to the presence of silver or copper in the culture medium. We report here on the development of an *in vivo* model to investigate the role of the SilABC proteins in the active transport of these heavy metal ions. The metal-sensitive *E. coli* strain GR17 was transformed with the *silABC* genes. Silver and copper tolerance of transformed bacteria was evaluated by the determination of minimal inhibitory concentration (MIC) values in a minimal Tris-glucose medium. A twofold increase in MIC values was observed when the three proteins were expressed demonstrating that the SilABC system transports efficiently silver and copper ions *in vivo*. In addition, we have demonstrated that the RND protein SilA alone is able to mediate partial resistance which is compatible with its cation/proton antiporter function.

2611-Pos**Effects of Various Inhibitors on Nitrate Uptake Mediated by *Bacillus* sp. GS2**

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Nitrate is one of the major components causing salt stress in Korean farm lands. In order to remove the excess amount of nitrate from the cultivating soils, soil bacteria having nitrate uptake activity were isolated. One of them showed a high capability of nitrate uptake. In the PCR analysis, 1373 bp of 16S rRNA gene were sequenced and compared to those of various microorganisms. The strain has identified as *Bacillus* sp. GS2. Growth of GS2 was not much facilitated by nitrate; however, high capacity of nitrate uptake was measured by removing 40 mM nitrate within 12 h. Nitrate transporter and nitrate reductase are useful enzymes to remove excess soil nitrate. GS2 showed high activity of nitrate reductase and the amount of nitrite formation was directly proportional to the amount of nitrate uptake. In order to characterize the bacterial nitrate uptake, the effect of chlorate was measured on the nitrate uptake activity of GS2 since chlorate was reported to inhibit nitrate transporter. While bacterial growth was not much inhibited by chlorate, the nitrate uptake was inhibited by 80% at the concentration of 50 mM chlorate. The effects of vanadate and phenylglyoxal (PGO) were measured on nitrate uptake. Both vanadate and chlorate showed similar patterns of inhibition on nitrate uptake and 50% inhibitions were obtained at 10-30 mM. PGO, an inhibitor of microbial nitrate transporter, completely inhibited the nitrate uptake at 1 mM. These results suggest that the nitrate reduction by GS2 is mediated by both membrane nitrate transporter and nitrate reductase in cytosol rather than by periplasmic enzyme.

2612-Pos**Physical and Functional Coupling of Electrogenic $\text{Na}^+/\text{HCO}_3^-$ Cotransport and Carbonic Anhydrases in the Myocardium**Alejandro Orlowski^{1,2}, Veronica C. de Giusti^{1,2}, Ernesto A. Aiello^{1,2}, Bernardo V. Alvarez^{1,2}.¹Centro de Invest. Cardiovasc., La PLATA, Argentina, ²FACULTAD DE CIENCIAS MEDICAS (UNLP), La Plata, Argentina.

The $\text{Na}^+/\text{HCO}_3^-$ cotransport (NBC) is an important sarcolemmal acid extruder in cardiac muscle. Functionally electroneutral (NBC3, 1 Na^+ :1 HCO_3^-) and electrogenic (NBC1 and NBC4, 1 Na^+ :2 HCO_3^-) forms of the transporter have been characterized in the heart. Mammalian cardiac muscle expresses membrane bound carbonic anhydrases (CA) IV (CAIV), IX (CAIX) and XIV (CAXIV), and cytosolic CAII. Association of CAII and CAIV, and NBC1, was previously demonstrated in the kidney and heterologous systems. NBC1

and CA physical and functional interaction was explored in the rat heart by co-immunoprecipitation and intracellular pH measurement (pH_i) experiments, respectively. CAII, CAIV, CAIX, and CAXIV were immunoprecipitated with anti-NBC1 antibody, using rat ventricular lysates. Conversely, non immune serum and irrelevant anti-glial fibrillar acidic protein antibodies failed to co-immunoprecipitate CAs with NBC1. NBC1 activity was investigated in isolated rat cardiomyocytes, using intracellular fluorescent measurements of BCECF-AM, to monitor pH_i . Cardiomyocyte membrane potential depolarizing pulses (MPDP) were applied by addition of 45 mM extracellular K^+ , to study NBC1 activity. After 10 minutes of MPDP a significant intracellular alkalinization was detected (0.17 ± 0.03 pH units; $n=6$, $P<0.05$). The alkalinization was fully cancelled with specific anti-NBC1 functionally inhibitory antibodies (0.02 ± 0.02 pH units; $n=5$), indicating activation of NBC1 isoform. Similarly, the NBC1-mediated increase of pH_i (0.17 ± 0.02 pH units; $n=11$, $P<0.05$), was inhibited with a poorly membrane-permeant CA inhibitor, benzolamide (100 μM , 0.09 ± 0.02 pH units; $n=6$, $P<0.05$), and a potent membrane-permeant CA inhibitor, ethoxylamide (100 μM , 0.05 ± 0.01 pH units; $n=6$, $P<0.05$), demonstrating a functional coupling between NBC1 cotransport and extracellular CAs, and NBC1 and intracellular CAs in the cardiac muscle, respectively. We demonstrated that the NBC1 $\text{Na}^+/\text{HCO}_3^-$ cotransport is functionally and physically coupled to both plasma membrane-anchored CAs and cytosolic CAII, forming a HCO_3^- transport metabolon in the myocardium.

2613-Pos**Role of $\text{Na}^+/\text{Ca}^{2+}$ Exchanger (NCX1) in Aldosterone-Induced Cardiac Remodeling**Takuya Iyoda¹, Satomi Kita¹, Shintaro Yamamoto¹, Issei Komuro², Akira Nishiyama³, Takahiro Iwamoto¹.¹Fukuoka University School of Medicine, Fukuoka, Japan, ²Chiba University Graduate School of Medicine, Chiba, Japan, ³Kagawa University School of Medicine, Kagawa, Japan.

Recent clinical and experimental studies have indicated the utility of eplerenone, a selective mineralocorticoid receptor antagonist, in cardiovascular and renal injuries. Actually, chronic treatment with aldosterone and salt can induce cardiac hypertrophy and renal injuries in experimental animals. However, its underlying mechanism is still unknown. To investigate whether the $\text{Na}^+/\text{Ca}^{2+}$ exchanger type 1 (NCX1) would be implicated in aldosterone-induced cardiac remodeling, we administered aldosterone (0.3 $\mu\text{g}/\text{h}$) and NaCl (1%) in NCX1 heterozygous mice (N1-KO), heart-specific transgenic mice overexpressing NCX1 (N1-Tg), and wild-type mice (WT). After 4 weeks of treatment, WT and N1-Tg showed a significant increase in heart weight/body weight (HW/BW) ratio with evidence of mild contractile dysfunction, whereas N1-KO maintained average HW/BW ratio and normal cardiac function. More importantly, NCX1 inhibitors (SEA0400, KB-R7943) attenuated aldosterone-induced cardiac hypertrophy and mild dysfunction when administered to WT for 4 weeks. On the other hand, prolonged treatment with eplerenone (200mg/kg/day) prevented cardiac hypertrophy that was spontaneously induced in N1-Tg. These results suggest that NCX1 participates in aldosterone-induced cardiac remodeling.

2614-Pos**Sodium-Dependent Inactivation of NCX1.3: Aortic Smooth Muscle Cells Versus Transfected CHO Cells**

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Despite the importance of Na/Ca exchange (NCX) activity in Ca homeostasis in blood vessels, there have been few detailed studies of the regulation of NCX activity in smooth muscle cells. Here we investigated the regulation of NCX by allosteric Ca activation and Na-dependent inactivation in rat aortic smooth muscle cells (ASMC) and compared the results with those of the smooth muscle isoform NCX1.3 expressed in CHO cells. Fura-2 loaded ASMC or transfected CHO cells were treated with ATP + thapsigargin to release Ca and prevent subsequent Ca re-accumulation in internal stores. Reverse exchange activity was initiated by applying 0.1mM Ca in Li-PSS or K-PSS. In both ASMC and transfected CHO cells treated with 1 mM ouabain, Ca uptake occurred following a 10-20sec lag period attributable to the positive feedback of allosteric Cai activation. To examine the Na-dependence of NCX activity, cells were treated with gramicidin and preincubated in 10-140 mM Na-PSS before activity was initiated by applying 0.1 mM CaCl_2 . In ASMC, NCX activity peaked at 20 mM Na but declined at higher Na concentrations; essentially no Ca uptake was seen at 140 mM Na. In contrast, robust activity was seen throughout that entire Na concentration range in transfected CHO cells. At 20 mM Na, Ca uptake in ASMC increased to a peak value and then declined sharply. After removing Ca with EGTA in 20 mM Na, subsequent pulses with 0.1 mM Ca revealed no activity until a 10 min recovery period had occurred. In contrast, NCX activity in the transfected CHO cells recovered